SEQUENCE LISTING

<110> Tibor Keler, et al.

<120> CELLS EXPRESSING ANTI-Fc RECEPTOR BINDING COMPONENTS

<130> MXI-099

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<150> 60/067,232
<151> 1997-12-02

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<170> PatentIn Ver. 2.0

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ctc tgg gtt cca ggt tcc act ggt gac tat cca tat gat gtt cca gat   157
Leu Trp Val Pro Gly Ser Thr Gly Asp Tyr Pro Tyr Asp Val Pro Asp
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tat gct ggg gcc cag ccg gcc aga tct gat atc cag ctg acc cag agc   205
Tyr Ala Gly Ala Gln Pro Ala Arg Ser Asp Ile Gln Leu Thr Gln Ser
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cca agc agc ctg agc gcc agc gtg ggt gac aga gtg acc atc acc tgt   253
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gcc tgg tac cag cag aag cca ggt aag gct cca aag ctg ctg atc tac   349
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gac atc gcc acc tac tac tgc cat caa tac ctc tcc tcg tgg acg ttc       493
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ggc caa ggg acc aag gtg gaa atc aag agc tct ggc ggt ggc ggc tcc       541
Gly Gln Gly Thr Lys Val Glu Ile Lys Ser Ser Gly Gly Gly Gly Ser
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gga ggt gga ggc agc gga ggg ggt gga tcc gag gtc caa ctg gtg gag       589
Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Glu Val Gln Leu Val Glu
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Ser Arg Asp Asn Ser Lys Asn Thr Leu Phe Leu Gln Met Asp Ser Leu
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aga ccc gaa gac acc ggg gtc tat ttt tgt gca aga ggc tac tat agg       877
Arg Pro Glu Asp Thr Gly Val Tyr Phe Cys Ala Arg Gly Tyr Tyr Arg
        255                 260                 265

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            Met Glu Thr Asp Thr Leu Leu Leu Trp Val Leu Leu
             1               5                   10

ctc tgg gtt cca ggt tcc act ggt gac tat cca tat gat gtt cca gat   157
Leu Trp Val Pro Gly Ser Thr Gly Asp Tyr Pro Tyr Asp Val Pro Asp
         15                  20                  25

tat gct ggg gcc cag ccg gcc aga tct gag atc cag ctg cag cag act   205
Tyr Ala Gly Ala Gln Pro Ala Arg Ser Glu Ile Gln Leu Gln Gln Thr
     30                  35                  40

gga cct gag ctg gtg aag cct ggg gct tca gtg aag ata tcc tgc aag   253
Gly Pro Glu Leu Val Lys Pro Gly Ala Ser Val Lys Ile Ser Cys Lys
 45                  50                  55                  60

gct tct ggt tat tca ttc act gac tac atc ata ttt tgg gtg aag cag   301
Ala Ser Gly Tyr Ser Phe Thr Asp Tyr Ile Ile Phe Trp Val Lys Gln
                 65                  70                  75

agc cat gga aag agc ctt gag tgg act gga aat aat aat cct tac tat   349
Ser His Gly Lys Ser Leu Glu Trp Thr Gly Asn Asn Asn Pro Tyr Tyr
             80                  85                  90

ggt agt act agc tac aat ctg aag ttc aag ggc aag gcc aca ttg act   397
Gly Ser Thr Ser Tyr Asn Leu Lys Phe Lys Gly Lys Ala Thr Leu Thr
             95                  100                 105

gta gac aaa tct tcc agc aca gcc tac atg cag ctc aac agt ctg aca   445
Val Asp Lys Ser Ser Ser Thr Ala Tyr Met Gln Leu Asn Ser Leu Thr
     110                 115                 120

tct gag gac tct gca gtc tat tac tgt gta aga gga gtt tat tac tac   493
Ser Glu Asp Ser Ala Val Tyr Tyr Cys Val Arg Gly Val Tyr Tyr Tyr
125                 130                 135                 140

ggt agt agc tac gag gcg ttt cct tac tgg ggc caa ggg act ctg gtc   541
Gly Ser Ser Tyr Glu Ala Phe Pro Tyr Trp Gly Gln Gly Thr Leu Val
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act gtc tct gca gga ggt ggc ggc tcc gga gga ggt ggc agc gga ggg     589
Thr Val Ser Ala Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly
            160                 165                 170

ggc gga tcc gat gtt gtg atg acc cag act cca ctc act ttg tcg att     637
Gly Gly Ser Asp Val Val Met Thr Gln Thr Pro Leu Thr Leu Ser Ile
            175                 180                 185

acc att gga caa cca gcc tcc atc tct tgc aag tca agt cag agc ctc     685
Thr Ile Gly Gln Pro Ala Ser Ile Ser Cys Lys Ser Ser Gln Ser Leu
            190                 195                 200

tta gat agt gat gga aag aca tat ttg aat tgg ttg tta cag agg cca     733
Leu Asp Ser Asp Gly Lys Thr Tyr Leu Asn Trp Leu Leu Gln Arg Pro
205                 210                 215                 220

ggc cag tct cca acg cgc cta atc tat ctg gtg tct aaa ctg gac tct     781
Gly Gln Ser Pro Thr Arg Leu Ile Tyr Leu Val Ser Lys Leu Asp Ser
            225                 230                 235

gga gtc cct gac agg ttc act ggc agt gga tca ggg aca gat ttc aca     829
Gly Val Pro Asp Arg Phe Thr Gly Ser Gly Ser Gly Thr Asp Phe Thr
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ctg aaa atc agc aga gtg gag gct gag gat ttg gga att tat tat tgc     877
Leu Lys Ile Ser Arg Val Glu Ala Glu Asp Leu Gly Ile Tyr Tyr Cys
            255                 260                 265

tgg caa ggt gca cat ttt cct cag acg ttc ggt gga ggc acc aag ctg     925
Trp Gln Gly Ala His Phe Pro Gln Thr Phe Gly Gly Gly Thr Lys Leu
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gaa atc aaa ccg cgg ctg cag gtc gac gaa caa aaa ctc atc tca gaa     973
Glu Ile Lys Pro Arg Leu Gln Val Asp Glu Gln Lys Leu Ile Ser Glu
285                 290                 295                 300

gag gat ctg aat gct gtg ggc cag gac acg cag gag gtc atc gtg gtg     1021
Glu Asp Leu Asn Ala Val Gly Gln Asp Thr Gln Glu Val Ile Val Val
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cca cac tcc ttg ccc ttt aag gtg gtg gtg atc tca gcc atc ctg gcc     1069
Pro His Ser Leu Pro Phe Lys Val Val Val Ile Ser Ala Ile Leu Ala
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ctg gtg gtg ctc acc atc atc tcc ctt atc atc ctc atc atg ctt tgg     1117
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